

First report of *Trichinella spiralis* from the white-eared (*Didelphis albiventris*) and the thick-tailed opossum (*Lutreolina crassicaudata*) in central Argentina

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Summary

Trichinellosis is a zoonotic disease caused by nematodes of the genus *Trichinella*. Humans, who are the final hosts, acquire the infection by eating raw or undercooked meat of different animal origin. *Trichinella spiralis* is an encapsulated species that infects mammals and is widely distributed in different continents. In Argentina, this parasite has been reported in the domestic cycle that includes pigs and synanthropic hosts (mainly rats and some carnivores). This is the first report of *T. spiralis* in the opossums *Didelphis albiventris* and *Lutreolina crassicaudata* in Argentina, and the first report in opossums in South America. In this survey, *Trichinella* larvae were detected by enzymatic digestion in three *D. albiventris* and one *L. crassicaudata* captured on pig and dairy farms located in the northeast of Buenos Aires province. The microscopic examination of the 32 larvae isolated presented the diagnostic characteristic of the genus *Trichinella*. Two larvae isolated from two *D. albiventris* and one from *L. crassicaudata* were identified as *T. spiralis* by nested multiplex PCR and confirmed by sequencing. Further research to determine the burdens of *T. spiralis* in opossums may contribute to a better understanding of the risk of *T. spiralis* transmission to the synanthropic populations.

Keywords: *Trichinella spiralis*; opossum; *Didelphis albiventris*; *Lutreolina crassicaudata*; Argentina

Introduction

Trichinellosis, the zoonotic disease also known as trichinosis or trichiniasis, is caused by nematodes belonging to the genus *Trichinella* (Murrell, 2007). The occurrence of trichinellosis in humans is strictly related to cultural food practices, including the consumption of raw or undercooked meat of different animal origin. *Trichinella* spp.

infection has been documented in both domestic (mainly pigs) and wild animals (Pozio, 2007). *T. spiralis*, widely distributed in different continents (Pozio, 2005), is the species involved in the domestic cycle that includes pigs and synanthropic hosts (like rats, marsupials and some carnivores). Humans accidentally acquire the infection by eating raw meat of infected pigs.

In Argentina, according to the current legislation, all slaughtered pigs in pork meat industries are subjected to HCl-pepsin digestion to avoid human infection. Nevertheless, pigs raised for own consumption and wild animals are not included in this legislation and are not controlled, thus representing the most important source of human infection. As a result, 5700 human cases have been reported in the last ten years (Bolpe, 2011). It has been proposed that the outdoor rearing of pigs is a major risk of transmission of *T. spiralis* because of the increased exposure to sylvatic and synanthropic hosts (mainly rats and some carnivores). *T. spiralis* can also be transmitted from pigs to synanthropic and sylvatic animals (Leiby *et al.*, 1990).

Trichinella spp. from sylvatic cycles have also caused human outbreaks as a result of the consumption of meat from puma (*Puma concolor*), armadillo (*Chaetophractus villus*) and wild boar (*Sus scrofa*) (Ribicich *et al.*, 2005). This parasite has also been detected in foxes (*Pseudalopex gracilis*) (Minoprio *et al.*, 1967). The recent description of the new species *T. patagoniensis* isolated from the puma in the Argentine Patagonia (Krivokapich *et al.*, 2012) makes it necessary to find out which wild animals are infected by each of these two *Trichinella* species as well as which are the areas of influence of the two parasites.

Marsupials are found in many different environments, from forests to domestic and peridomestic habitats like orchards, henhouses and poultry farms (Cabrera & Yepes, 1960; Hunsaker II, 1977; Contreras, 1983; Gómez Villa-

fañe *et al.*, 2004). Since they have been described as reservoirs of numerous diseases (Potkay, 1977; Schweigmann *et al.*, 1999; Gomes *et al.*, 2003; Bodini Santiago *et al.*, 2007; Pérez Carusi *et al.*, 2009), some marsupials are considered a link between wild and domestic habitats for the transmission of zoonotic diseases (Gómez Villafañe *et al.*, 2004; Pérez Carusi *et al.*, 2009).

In environments related to the human population, opossums of the genus *Didelphis* are associated with garbage produced by humans, indicating the close relation between these animals and humans (Pérez Carusi *et al.*, 2009). *Didelphis albiventris*, the white-eared opossum, is one of the six species of the genus. It is widely distributed in the American continent, from eastern North America throughout eastern South America, from northeastern Brazil through Paraguay, Uruguay and southern Argentina, up to 47 °S (Cabrera & Yepes, 1960; Cerqueira, 1985; Navone & Suriano, 1992; Lemos & Cerqueira, 2002). Although *D. albiventris* is mainly insectivorous, it may adopt omnivorous feeding habits, including fruits and vertebrates in its diet (Cabrera & Yepes, 1960; Navone & Suriano, 1992; Cáceres, 2002), thus appearing to be opportunistic (Aléssio *et al.*, 2005).

Lutreolina crassicaudata, the thick-tailed opossum, which is the only living species of the genus, also has a wide distribution, occurring in southern Brazil, Argentina, Uruguay, Paraguay and part of Bolivia, as well as in areas of northern South America, such as Colombia, Venezuela and Guyana (Redford & Eisenberg, 1992; Flores *et al.*, 2007). Although *L. crassicaudata* is less studied than *D. albiventris* and other didelphid marsupial species (Santori *et al.*, 2005), it is known that this marsupial has omnivorous habits (Cáceres *et al.*, 2002; Muschetto *et al.*, 2011). *Monodelphis dimidiata*, the southern short-tailed opossum, and *Gracilinanus agilis*, the gracile mouse opossum, are two other opossums that occur in our study area but in very low abundances.

The aim of this work is to report the first findings of *Trichinella* in the opossums *D. albiventris* and *L. crassicaudata* on intensive animal breeding farms in central Argentina.

Materials and methods

Study area

The fieldwork was carried out in the localities of Marcos Paz, Las Heras, Exaltación de la Cruz and San Andrés de Giles located in the northeast of Buenos Aires province, Argentina. The climate is temperate with a mean annual temperature of 17.4 °C (IGM, 1998). The original pastures have been replaced by grain crops and natural or implanted pastures for livestock breeding. The intensive poultry, porcine and bovine cattle farming is also characteristic of this rural area. Trichinellosis is the parasitary disease with highest incidence in Buenos Aires province, with 2387 human cases reported from 2000 to 2010. This is half of the total cases for all the country.

Opossum survey and larvae identification

This study was carried out on porcine and dairy farms as a part of a larger survey where the ecology and parasitology of synanthropic mammals were studied. Trapping campaigns were conducted on ten pig farms and eight dairy farms from December 2008 to November 2011. Each of these farms was sampled four times along one year (once each season). Opossums were captured using cage live traps (15 x 16 x 31 cm) set in lines in the perimeter of pig sheds or dairies, houses, machinery sheds, food storage sheds or silos and in vegetated areas (including lawn areas, stream weedy borders and weedy borders below fences). The distance between neighboring traps along the lines was 5 to 10 m. Traps were active for three consecutive nights and baited with beef and carrot. The opossums captured were identified to species level. Animals were euthanized with an overdose of ketamine-acepromazine according to Pérez Carusi *et al.* (2009). The tongue, diaphragm, intercostals and leg muscles were removed and preserved at -30 to 0 °C. A pool of the total muscle samples from each animal was analyzed by artificial digestion in 1 % HCl, 1 % pepsin (Gamble *et al.*, 2000). All larvae isolated were morphologically identified using an inverted microscope. Genomic DNA from all single muscle larvae was extracted and analyzed by nested-multiplex PCR according to Zarlenga *et al.* (1999). The expansion segment five (ESV) was amplified. Muscle larvae of the *Trichinella* spp. reference isolates *T. spiralis* (ISS599) and *T. patagoniensis* (ISS2311) were used as controls. Forward and reverse sequencing was carried out on PCR products by a capillary DNA sequencer ABI 3730xl DNA analyzer (Applied Biosystems, Macrogen Inc., Seoul, South Korea). Nucleotide sequences were aligned with published sequences for *T. spiralis* (Mallatt & Winchell, 2002) using the software of the European Bioinformatics Institute website (<http://www.ebi.ac.uk/Tools/clustalw2/>).

Results

A total of 61 opossums (41 *D. albiventris* and 20 *L. crassicaudata*) were captured during the study period, with a total trapping effort of 7026 trap-nights. *D. albiventris* was captured on four dairy farms and six porcine farms, while *L. crassicaudata* was captured on three dairy farms and one porcine farm.

Muscle samples of a total of 59 opossums were processed (mean weight = 9.01 g, standard deviation = 6.11 g, minimum = 1.45 g, maximum = 24.00 g), while samples of two opossums were lost. We found 32 *Trichinella* spp. larvae in three *D. albiventris* (with 0.67, 2.81 and 1.36 larvae per gram), and one *L. crassicaudata* (with 0.26 larvae per gram). Two of the infected opossums were captured on a stream border of one dairy farm (one *D. albiventris* and the *L. crassicaudata*), while the other two *D. albiventris* were captured on a food shed of another dairy farm and on a breeding shed of a porcine farm. The examination of the 32 larvae under microscope showed the presence of the stichosome, which is a diagnostic characteristic of the genus

Trichinella. The other morphological characteristics observed were also consistent with this genus. Additionally, one larva isolated from a *D. albiventris* could not be identified, but it did not have the typical morphological characteristics of the genus *Trichinella*.

The genomic DNA extraction was successful for 15 out of the 32 larvae found, and in three of these 15 the ESV was amplified successfully. Two larvae isolated from two *D. albiventris* and only one from *L. crassicaudata* were identified as *T. spiralis* by nested multiplex PCR, generating the 173-bp fragment consistent with the pattern for *T. spiralis* (Fig. 1). Two isolates, one from *D. albiventris* and the other from *L. crassicaudata*, were also confirmed as *T. spiralis* by sequencing. CLUSTALW2 alignments showed sequences identical to those deposited by Mallatt & Winchell (Mallatt & Winchell, 2002) at GenBank (accession number AF342803.1).

36 *D. albiventris* studied. Less is known about *L. crassicaudata*. In central Argentina, this species has been found parasitized by the helminths *Turgida turgida*, *Pterygodermatites kozeki*, *Aspidodera raillieti*, by the nematodes *Mathevotaenia* sp., and by the acanthocephalans *Hamanniella microcephala* (Navone *et al.*, 1991). In Florianópolis, Brazil, *L. crassicaudata* has been found parasitized by the protozoan *Tetratrichomonas didelphidis* (Tasca *et al.*, 2001).

The number of opossums captured in this study was low to estimate the prevalence in the populations. More individuals of *D. albiventris* and *L. crassicaudata* in a longer period of time should thus be analyzed to determine the role of these opossums as reservoirs or sporadic hosts of *T. spiralis* in the study area. Additional studies about the role of these opossums for pig infection with *T. spiralis* are also needed. However, the fact that we detected infected ani-

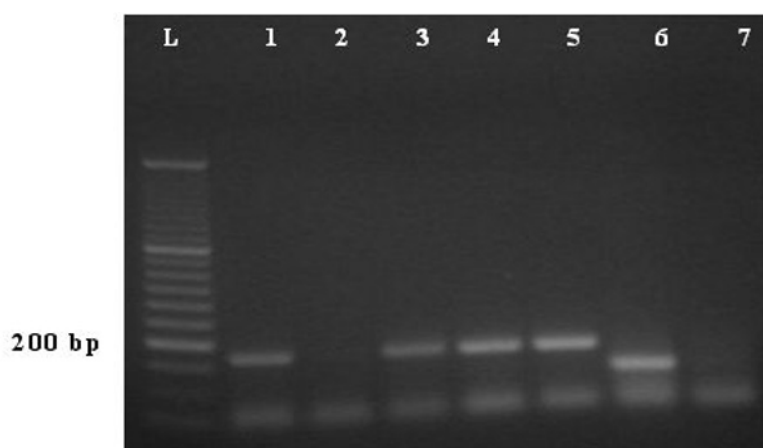


Fig. 1. Agarose gel separation of nested multiplex PCR products from: (1) single larva collected from one *D. albiventris*, (2) single larva collected from one *D. albiventris* that showed no features of the genus *Trichinella*, (3) single larva collected from another *D. albiventris*, (4) single larva collected from one *L. crassicaudata*, (5) *T. spiralis* reference strain (ISS599), (6) *T. patagoniensis* reference strain (ISS2311), (7) negative control, (L) 50-bp ladder

Discussion

Knowledge about pathogens of *L. crassicaudata* and *D. albiventris* is scarce and there are no previous reports of *T. spiralis* infection, being our findings the first report of *T. spiralis* in these species and the first report in opossums in South America. In Argentina, some studies looked for this infection in *D. albiventris* with negative results (Gómez Villafañe *et al.*, 2004; Pérez Carusi *et al.*, 2009; Ribicich *et al.*, 2010). In contrast, in the USA, two opossums, *D. marsupialis* and *D. virginiana*, have been previously reported to be infected with *Trichinella* spp. (Pozio, 2005).

As mentioned, in Argentina, some opossums are considered a potential link between wild and domestic habitats for the transmission of zoonotic diseases (Gómez Villafañe *et al.*, 2004; Pérez Carusi *et al.*, 2009). Pérez Carusi *et al.* (2009) conducted a screening of different pathogens in *D. albiventris* in a rural area of central Argentina and found *Leptospira interrogans* and *Salmonella enterica* in 13 % and 4 % of the 179 individuals tested, respectively, but found no individuals infected by *T. spiralis*. Moreover, Ribicich *et al.* (2010) reported no *Trichinella* spp. in other

imals suggests that these opossums could participate in the transmission cycle of *T. spiralis*. The infected opossums were all captured on pig and dairy farms and one *D. albiventris* was found inside a food storage shed of a dairy farm with high human activity. Since these opossums have home ranges that exceeded the farm limits, with moving areas around the nesting place of 7.04 ha for *D. albiventris* (Sanches Quadros Altomare *et al.*, 2012) and 0.95 ha for *L. crassicaudata* (Cajal, 1981), they could use the surrounding rural habitats of the farms. Also, individuals of Norway rat (*Rattus norvegicus*), black rat (*R. rattus*), and house mouse (*Mus musculus*) have been previously found infected with *T. spiralis* on the farms studied in the present survey (Lovera R, unpublished data). For all these reasons, we agree with the idea that these opossum species could act as a link between wild and domestic habitats for the transmission of zoonotic diseases as Trichinellosis.

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